

Dimensionality reduction and clustering

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 An abbreviated version of this protocol was published in eLIFE in Jan 2021

A transcriptomic taxonomy of *Drosophila* circadian neurons around the clock

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Detailed protocol

The code we used for dimensionality reduction, clustering and rhythmic gene expression analysis has been deposited in GitHub (https://github.com/rosbash-lab/scRNA_seq_clock_neurons).

How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Ma, D. , Przybylski, D. , Abruzzi, K. , Schlichting, M. , Li, Q. , Long, X. and Rosbash, M. (2021). Dimensionality reduction and clustering. Bio-protocol Preprint. bio-protocol.org/prep833.
2. Ma, D., Przybylski, D., Abruzzi, K. C., Schlichting, M., Li, Q., Long, X. and Rosbash, M.(2021). A transcriptomic taxonomy of *Drosophila* circadian neurons around the clock. eLIFE. DOI: [10.7554/eLife.63056](https://doi.org/10.7554/eLife.63056)

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